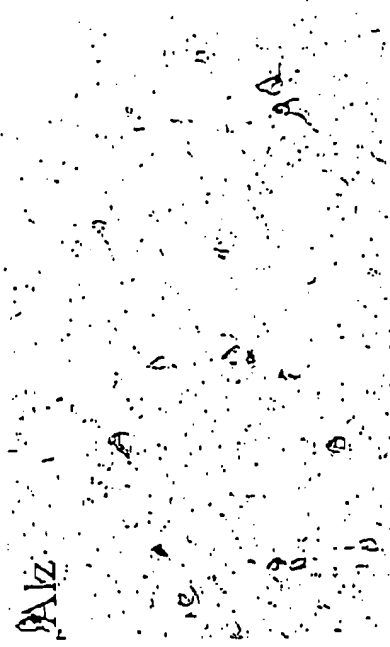


Figure 1

AD2

AD46



T

T

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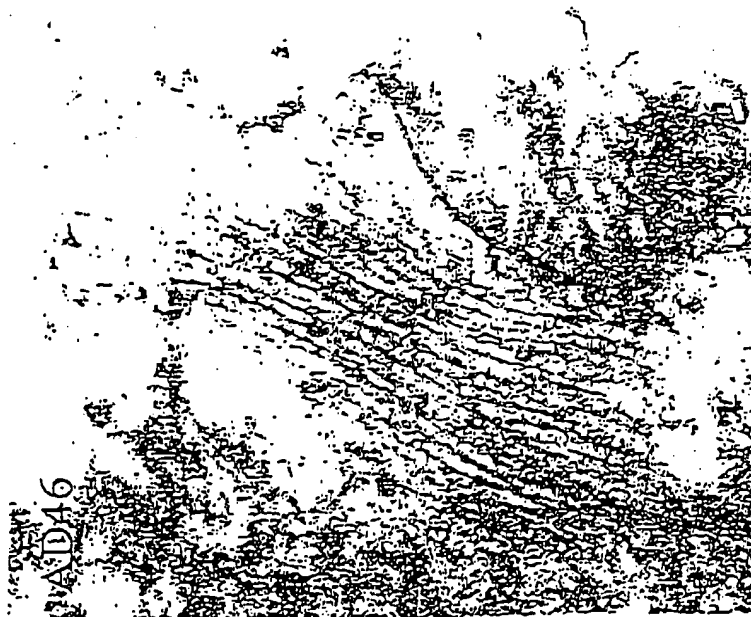
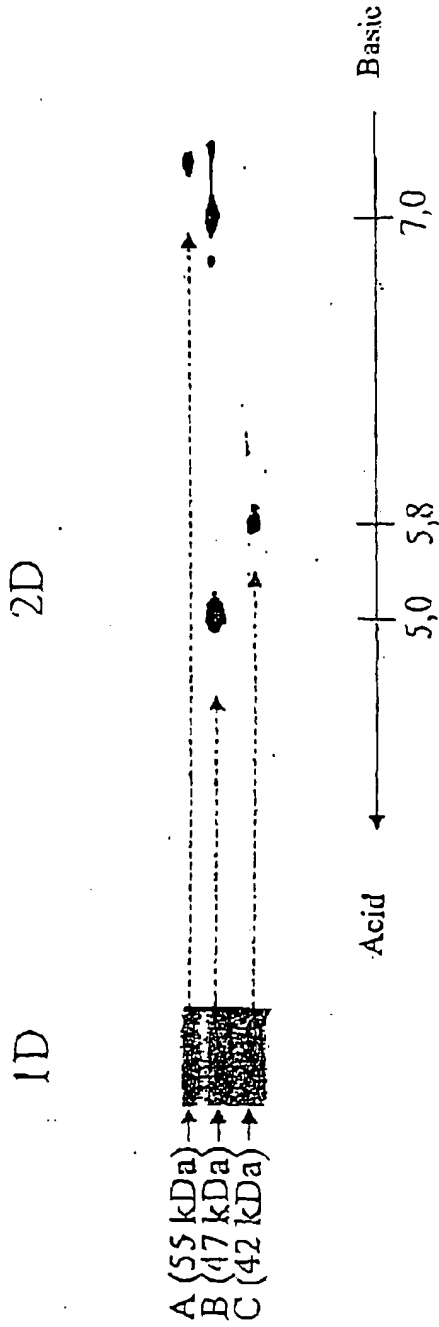


Figure 2

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Figure 3



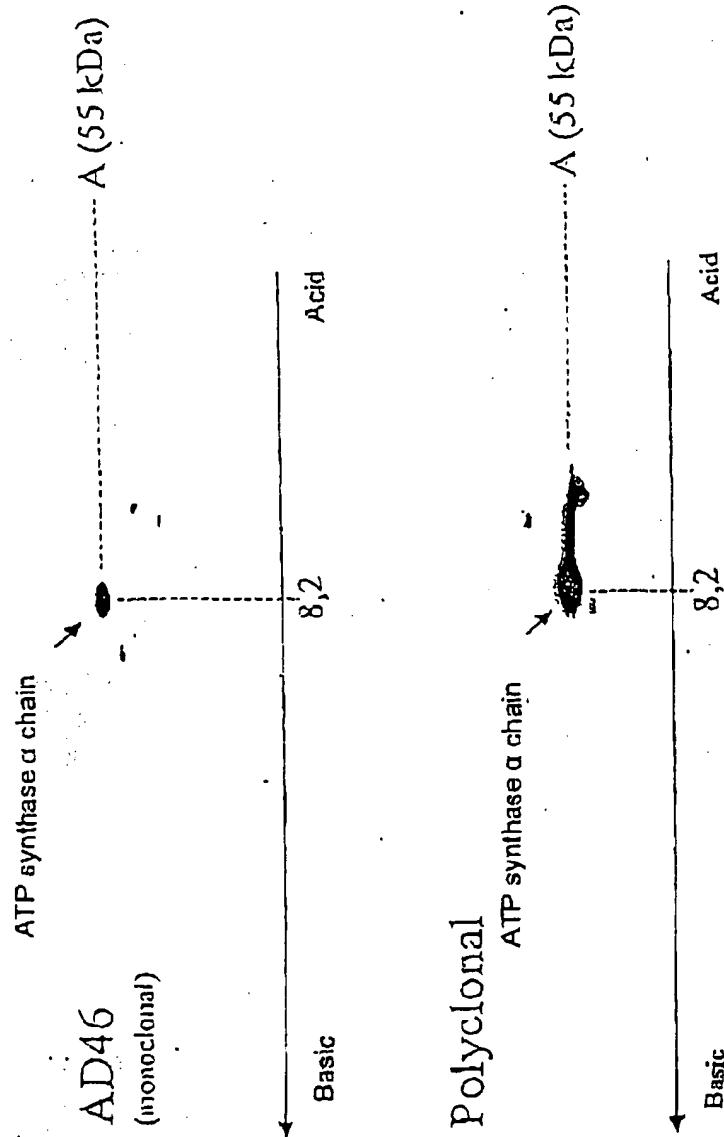
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Figure 4

MM of the fragments of protein A					
815.459	892.494	1026.589	1149.614	1287.703	1438.872
1553.783	1551.829	1824.928	1867.8	1683.797	1927.327
					2120.029
Identification score	376				
% of fragments identified	75				
Species	human				
Identification number	P06576				
Name	ATP synthase $\alpha$ chain (EC 3.5.1.34)				
MM of the fragments of protein B					
848.299	880.299	1160.199	1570.499	1583.499	1805.199
1938.899	2366.499	2482.199	2578.699	2702	1858.899
Identification score	164.3				
% of fragments identified	60				
Species	human				
Identification number	P00104				
Name	Gamma Enolase (EC 4.2.1.11)				
MM of the fragments of protein C					
784.899	945.189	975.999	1132.299	1198.499	1516.099
1953.299	2214.599				1790.689
Identification score	178.5				
% of fragments identified	100				
Species	human				
Identification number	P15475				
Name	Cytoplasmic actin				

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Figure 5



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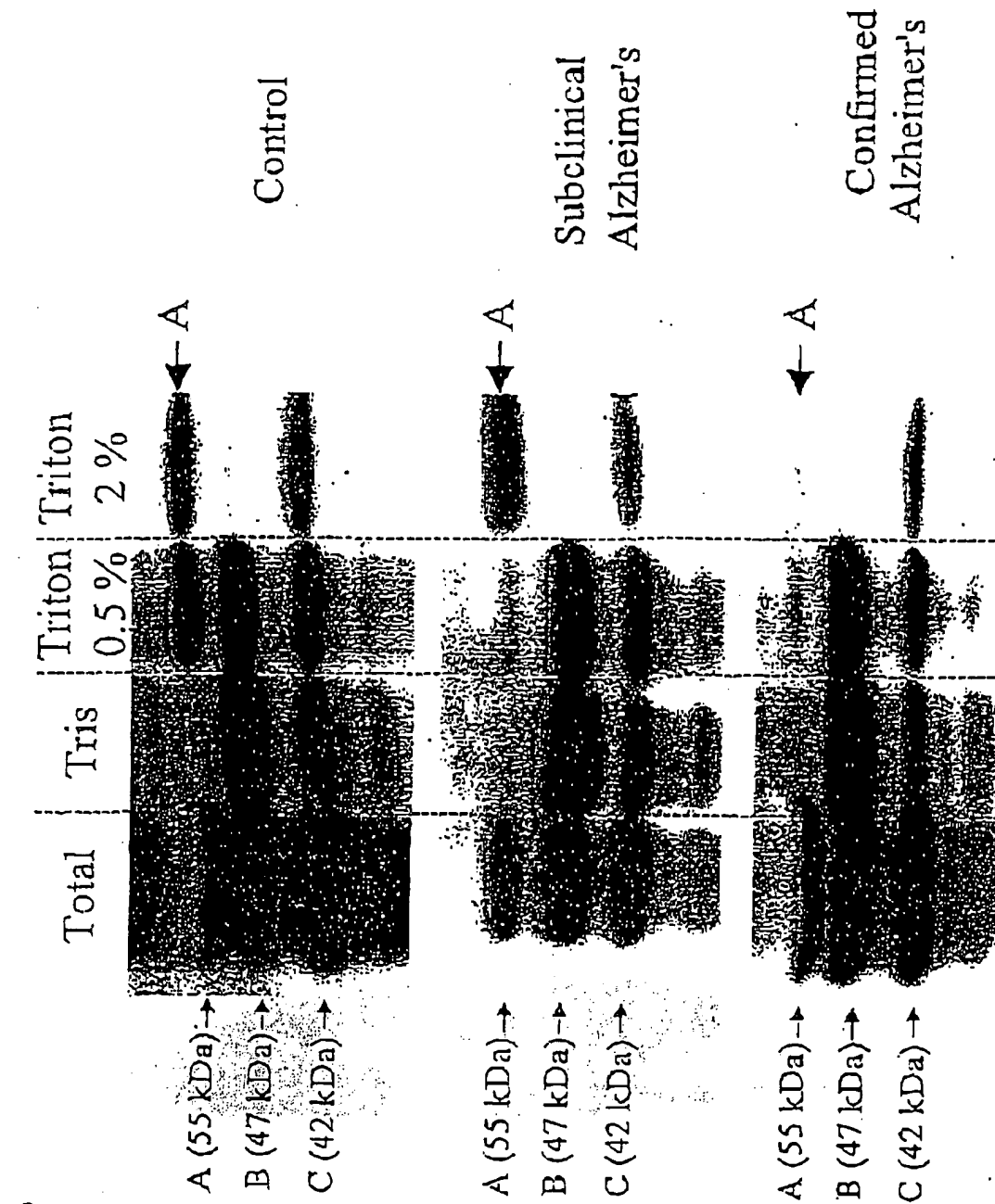


Figure 6

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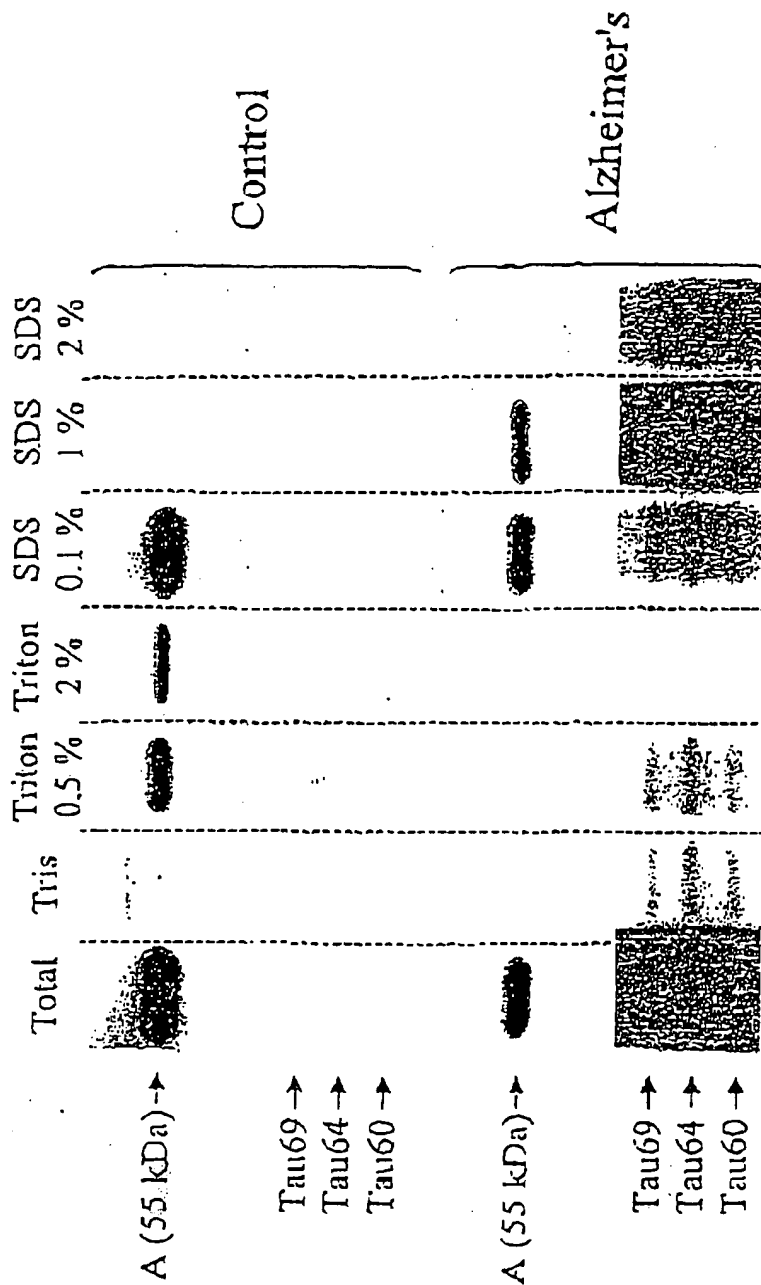


Figure 7.

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**A**

Control

20K 100K 100K  
P S P

Total

Polyclonal  
Anti-ATP synthase  
α chain

Polyclonal  
Anti-ATP synthase  
α chain

**B**

Immunoprecipitation

AD46 ATPΣ  
TauCl AD2

Marking

AD46 ATPΣ  
TauCl AD2

Alz

20K 100K 100K  
P S P

Total

Alz

Polyclonal  
Anti-ATP synthase  
α chain

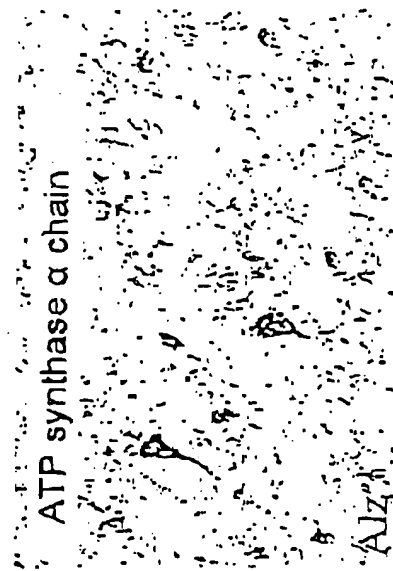
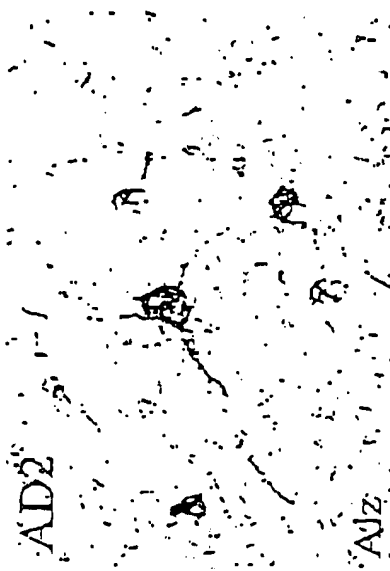
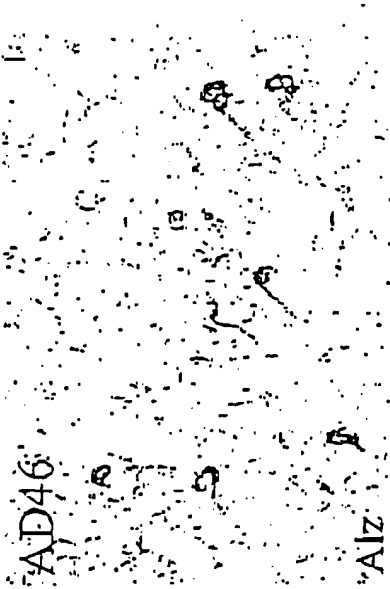
169  
164  
160

169  
164  
160

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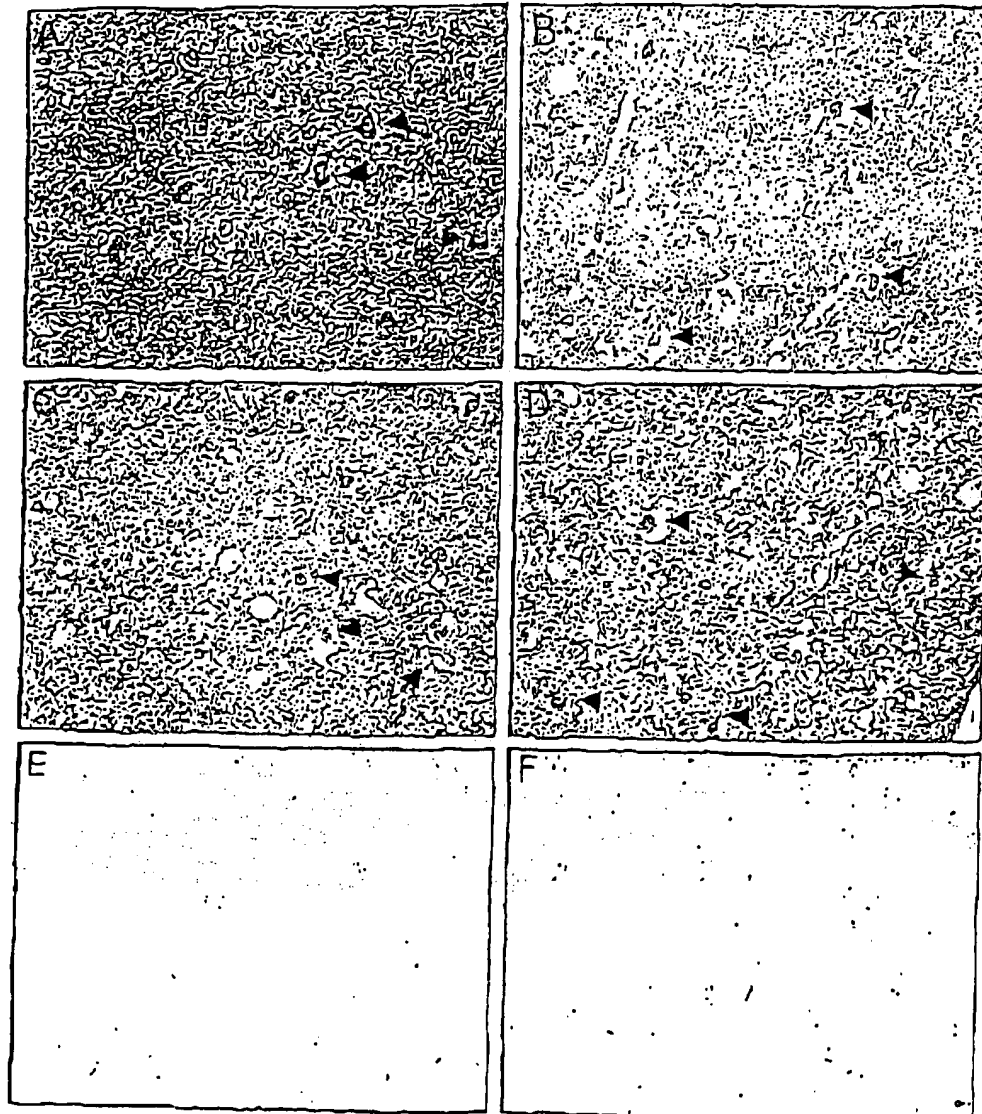


Figure 9



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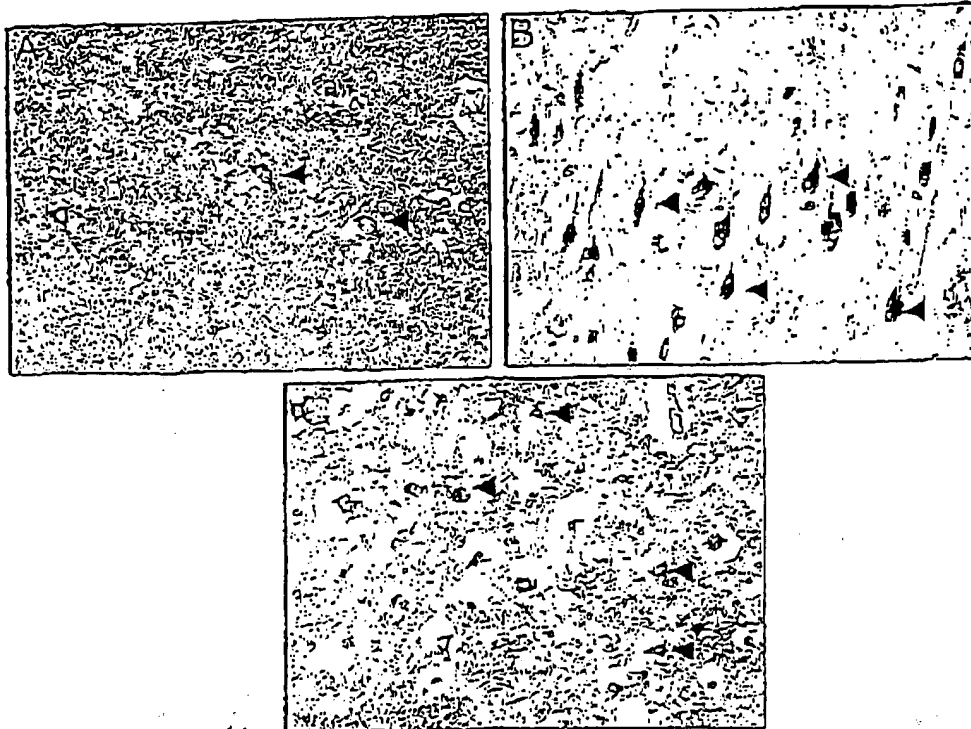
Figure 10:



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Figure 11:



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